

SEQUENCE LISTING

<110> Brady, William
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<120> LIGAND FOR CD28 RECEPTOR ON B CELLS AND METHODS

<130> ON0065C Div 1/30436.11US06

<140> 09/666,267
<141> 2000-09-21

<150> 07/498,949
<151> 1990-03-26

<150> 07/547,980
<151> 1990-07-02

<150> 07/722,101
<151> 1991-06-27

<150> 08/219,200
<151> 1994-03-29

<150> 08/459,766
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<170> PatentIn version 3.1

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<301> Gordon J. Freeman, Arnold S. Freedman, Jeffrey M. Segil, Grace Lee, James F. Whitman and Lee M. Nadler
<302> B7, A NEW MEMBER OF THE Ig SUPERFAMILY WITH UNIQUE EXPRESSION ON ACTIVATED AND NEOPLASTIC B CELLS
<303> J. Immunol.
<304> 143
<305> 8
<306> 2714 TO 2722
<307> 1989-10-15
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<313> (1)..(216)

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